

SEQUENCE LISTING

- (1) GENERAL INFORMATION:
 - (i) APPLICANT: Gunzburg, Walter H. Winder, David Saller, Robert M.
 - (ii) TITLE OF INVENTION: VECTORS CARRYING THERAPEUTIC GENES ENCODING ANTIMICROBIAL PEPTIDES FOR GENE THERAPY
 - (iii) NUMBER OF SEQUENCES: 12
 - (iv) CORRESPONDENCE ADDRESS:
 - (A) ADDRESSEE: HAMILTON, BROOK, SMITH & REYNOLDS, P.C.
 - (B) STREET: TWO MILITIA DRIVE
 - (C) CITY: LEXINGTON
 - (D) STATE: MASSACHUSETTS
 - (E) COUNTRY: USA (F) ZIP: 02173
 - (v) COMPUTER READABLE FORM:
 - (A) MEDIUM TYPE: Floppy disk
 - (B) COMPUTER: IBM PC compatible
 - (C) OPERATING SYSTEM: PC-DOS/MS-DOS
 - (D) SOFTWARE: PatentIn Release #1.0, Version #1.30
 - (vi) CURRENT APPLICATION DATA:
 - (A) APPLICATION NUMBER: US 08/999,690
 - (B) FILING DATE: 08-SEP-1997
 - (C) CLASSIFICATION:
 - (vii) PRIOR APPLICATION DATA:
 - (A) APPLICATION NUMBER: PCT/EP96/01001
 - (B) FILING DATE: 08-MAR-1996
 - (vii) PRIOR APPLICATION DATA:
 - (A) APPLICATION NUMBER: DK 0243/95
 - (B) FILING DATE: 09-MAR-1995
 - (viii) ATTORNEY/AGENT INFORMATION:
 - (A) NAME: Brook, David E.
 - (B) REGISTRATION NUMBER: 22,592
 - (C) REFERENCE/DOCKET NUMBER: GSF97-03A
 - (ix) TELECOMMUNICATION INFORMATION:
 - (A) TELEPHONE: (781) 861-6240
 - (B) TELEFAX: (781) 861-9540

- (2) INFORMATION FOR SEQ ID NO:1:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 26 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: unknown
 - (D) TOPOLOGY: unknown
 - (ii) MOLECULE TYPE: protein
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:

Gly Ile Gly Ala Val Leu Lys Val Leu Thr Thr Gly Leu Pro Ala Leu 1 5 10 15

Ile Ser Trp Ile Lys Arg Lys Arg Gln Gln 20 25

- (2) INFORMATION FOR SEQ ID NO:2:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 26 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: unknown
 - (D) TOPOLOGY: unknown
 - (ii) MOLECULE TYPE: protein
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:

Gly Ile Gly Ala Val Leu Lys Val Leu Thr Thr Gly Leu Pro Ala Leu 1 5 10 15

Ile Ser Trp Ile Gly Gly Gly Gly Gly Gly 20 25

- (2) INFORMATION FOR SEQ ID NO:3:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 25 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: other nucleic acid
 (A) DESCRIPTION: /desc = "DNA"
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:

TATGACGTCT CGTTAGAACG CGGCT

(2)	INFORMATION FOR SEQ ID NO:4:	
	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
	<pre>(ii) MOLECULE TYPE: other nucleic acid (A) DESCRIPTION: /desc = "DNA"</pre>	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:	
GGC	AGATCTT AAATGTATCA TGCAAT	26
(2)	INFORMATION FOR SEQ ID NO:5:	
	(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 6 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	<pre>(ii) MOLECULE TYPE: other nucleic acid (A) DESCRIPTION: /desc = "DNA"</pre>	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:	
GAC		6
(2)	INFORMATION FOR SEQ ID NO:6:	
	(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 6 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	<pre>(ii) MOLECULE TYPE: other nucleic acid (A) DESCRIPTION: /desc = "DNA"</pre>	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:	
AGAT	TCT	6
(2)	INFORMATION FOR SEQ ID NO:7:	
	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 28 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	

<pre>(ii) MOLECULE TYPE: other nucleic acid</pre>	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:7:	
ATAGACGTCA AGGAAGGAAG CGATCGGA	28
(2) INFORMATION FOR SEQ ID NO:8:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 28 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
<pre>(ii) MOLECULE TYPE: other nucleic acid (A) DESCRIPTION: /desc = "DNA"</pre>	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:8:	
TATGGATCCA ACCCTGTTGC CTCTTACG	28
(2) INFORMATION FOR SEQ ID NO:9:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 35 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
<pre>(ii) MOLECULE TYPE: other nucleic acid</pre>	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:9:	
TCTTACATCT ATGCGGGAAT TGGAGCAGTT CTGAA	35
(2) INFORMATION FOR SEQ ID NO:10:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 35 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
<pre>(ii) MOLECULE TYPE: other nucleic acid (A) DESCRIPTION: /desc = "DNA"</pre>	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:10:	
AACTGCTCCA ATTCCCGCAT AGATGTAAGA AATGT	35

(2) INFORMATION FOR SEQ ID NO:11:

(i) SEQUENCE CHARACTERISTICS:

	(A) LENGTH: 6 base pairs(B) TYPE: nucleic acid(C) STRANDEDNESS: single(D) TOPOLOGY: linear	
(<pre>(ii) MOLECULE TYPE: other nucleic acid (A) DESCRIPTION: /desc = "DNA"</pre>	
•	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:11:	
GACGT	TC	6
(2)	INFORMATION FOR SEQ ID NO:12:	
	(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 6 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
•	<pre>(ii) MOLECULE TYPE: other nucleic acid (A) DESCRIPTION: /desc = "DNA"</pre>	
•	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:12:	
GGATO	CC	6